LabGym: **quantification of user-defined animal behaviors using learning-based holistic**

assessment

Yujia Hu[1,*], Carrie R. Ferrario[2], Alexander D. Maitland[2], Rita B. Ionides[2], Anjesh Ghimire[3],

Brendon Watson[3], Kenichi Iwasaki[1], Hope White[1], Yitao Xi[1], Jie Zhou[4,*], Bing Ye[1,*]

[1] Life Sciences Institute and Department of Cell and Developmental Biology, University of

Michigan, Ann Arbor, MI 48109, USA

[2] Department of Pharmacology and Psychology Department (Biopsychology), University of

Michigan, Ann Arbor, MI 48109, USA

[3] Department of Psychiatry, University of Michigan, Ann Arbor, MI 48109, USA

[4] Department of Computer Science, Northern Illinois University, DeKalb, IL 60115, USA

* Correspondence: Yujia Hu (henryhu@umich.edu), Jie Zhou (jzhou@niu.edu) and Bing Ye

(bingye@umich.edu)

ABSTRACT

Quantifying animal behavior is important for many branches of biological research. Current

computational tools for behavioral quantification typically rely on a few pre-defined, simplified

features to identify a behavior. However, such an approach restricts the information used and

the tool's applicability to a limited range of behavior types or species. Here we report a new tool,

LabGym, for quantifying animal behaviors without such limitations. Combining a novel approach

for effective evaluation of animal motion with customizable convolutional recurrent networks for

capturing spatiotemporal details, *LabGym* provides holistic behavioral assessment and

accurately identify user-defined animal behaviors without restrictions on behavior types or

animal species. It then provides quantitative measurements of each behavior, which quantify the

behavior intensity and the body kinematics during the behavior. *LabGym* requires neither any

intermediate step for processing features that causes information loss nor programming

knowledge from users for post-hoc analysis. It tracks multiple animals simultaneously in various

experimental settings for high-throughput and versatile analysis. It also provides users a way to

generate visualizable behavioral datasets that are valuable resources for the research

community. We demonstrate its efficacy in capturing subtle behavioral changes in animals

ranging from soft-bodied invertebrates to mammals.

INTRODUCTION

Quantitative measurements of animal behavior are important for many branches of biology and

biomedical research (Altimus et al., 2020; Datta et al., 2019; Egnor and Branson, 2016; Pereira

et al., 2020). Identifying a behavior is a prerequisite to quantifying it. Current computational tools

for behavioral quantification largely rely on tracking a set of pre-defined, simplified features to

identify a behavior (Berman et al., 2014; Denisov et al., 2013; Hsu and Yttri, 2021; Kabra et al.,

2013; Klibaite et al., 2017; Klibaite and Shaevitz, 2020; Nilsson et al., 2020; Pereira et al., 2019;

Segalin et al., 2021; Wiltschko et al., 2015). For example, JAABA uses pre-defined features

(e.g., instantaneous speed, positions, and areas) that are derived from the outlines of the

animals' bodies for identifying a set of behaviors in mice and flies (Kabra *et al.*, 2013). Similarly,

B-SOiD uses body part positions computed by DeepLabCut (Mathis et al., 2018) to generate

speed, angles, and body lengths for identifying animal behaviors (Hsu and Yttri, 2021). MARS

uses the postural features of two mice to classify social behaviors in mice (Segalin *et al.*, 2021).

MotionMapper also uses postural features of adult *Drosophila* to identify behaviors in flies

(Berman *et al.*, 2014). In *Drosophila* larvae, SALAM/LARA evaluates several specific contour

features for behavior detection (Denisov *et al.*, 2013).

While tools that use pre-defined, simplified features have greatly advanced behavioral

analyses, they have limitations. Firstly, the features need to be defined *a priori* for identifying

certain behavior types, which may fail to generalize to other behavior types. For example,

motion speed can be used to identify locomotion but is useless for distinguishing facial

grooming from body grooming during which the animal is immobile. Animals can also display

new behaviors in response to experimental manipulations, such as drug exposure (Ellinwood

and Balster, 1974; Ferrario et al., 2005). Such qualitatively different behaviors might not be

easily captured by pre-defined features. Moreover, the same behavior may be recorded from

different camera angles. A tool with features that are pre-defined for side views might not work

for videos captured in top-view. Secondly, feature-based approaches assume that only the

features selected by the designer are key to identifying a given behavior. However, the limited

number of pre-defined features typically underrepresent the complete information needed to

capture the complexity of animal behavior; and defining a large set of features that encompass

every aspect of a behavior *a priori* is not only computationally inefficient but also might be

impossible. Finally, a feature is typically a simplified summary of one aspect of the behavior.

Information loss might occur in processing the feature, which makes the behavior identification

less accurate. For example, speed derived from the distance of the animals' centers of mass is

a common feature that is used in feature-based tools, but the information of animal body shape

is lost in such simplification.

Recent advancements in deep learning offer new approaches for behavior analysis. For

example, DeepLabCut and LEAP use deep convolutional neural networks (Fukushima, 1980;

LeCun and Bengio, 1995) to enable accurate tracking of animal body parts for pose estimation

(Mathis *et al.*, 2018; Pereira *et al.*, 2019). These tools track body positional in space across time,

but on their own do not identify or quantify any behavior. As a result, additional post-hoc tools,

such as B-SOiD (Hsu and Yttri, 2021), are required to identify and quantify a behavior based on

the outputs of the tracking tools. Other deep-learning-based tools such as DeepEthogram

(Bohnslav et al., 2021) and SIPEC (Marks et al., 2022) focus on behavior classification but do

not provide quantitative measurements of the behavior intensity or the body kinematics during a

behavior, which are key to behavioral analysis. Moreover, they are computationally expensive

and require powerful graphics processing units (GPUs) to run.

Here, we report a new tool, *LabGym*, for quantifying user-defined animal behaviors

without restrictions on behavior types or animal species, or the need to apply pre-defined or

simplified features for behavior identification. When people manually analyze behaviors in a

video, they typically ignore irrelevant information (such as static background) and make holistic

observations of the behaving animal by considering both spatiotemporal changes and the

overall movement pattern to categorize the behavior. Inspired by this human cognitive process,

we designed *LabGym* to ignore static backgrounds, track multiple animals, accurately

categorize behavior by holistic assessment, and then calculate quantitative measurements of

each behavior. The technical innovations in *LabGym* include customizable combinations of

neural networks for effective assessment of the spatiotemporal information in a behavior

episode of user-defined duration, a new approach that we designed for efficiently evaluating the

animal's movement pattern during a behavior, and a new method for effective background

subtraction. These innovations have made *LabGym* run easily on a common laptop computer,

while most current deep-learning-based tools for behavioral analysis require demanding

computational capabilities (Bohnslav *et al.*, 2021; Marks *et al.*, 2022; Mathis *et al.*, 2018).

Moreover, *LabGym* provides users a way to generate visualizable behavioral datasets

that can be shared across the research community as ground truth (i.e., the data with assigned

true labels) (LeCun et al., 2015; Roh et al., 2021; Schmidhuber, 2015) or for benchmarking

different algorithms. Furthermore, *LabGym* includes real-time visualizations of behavioral

categorizations and a graphical user interface (GUI) containing flexible options for

experimenters to use without the need of writing computer code.

1    We show that *LabGym* precisely captures subtle changes in user-defined behaviors in

2 animals ranging from soft-bodied invertebrates to mammals. We further demonstrate its

3 strength by comparing it with the existing standards.

RESULTS

The *LabGym* pipeline

LabGym (https://github.com/umyelab/LabGym) was designed to mimic human assessment of
animal behaviors. Our goals were to design software that efficiently and accurately identifies all
the user-defined behavioral types (i.e., "categories") for each animal in an experimental session,
and that provides quantitative measurements of each category.

To accomplish this, *LabGym* first uses a new method to effectively remove background
in each video frame and tracks each animal within the video (Figure 1A). At each frame of the
video, a module termed the 'Categorizer' determines which behavior an animal performs during
a time window that spans from the current frame back to a user-defined number of frames prior
('time window'). To achieve this, two types of data are extracted at each video frame for each
animal: a) a background-free animation ('animation') that spans the specified time window, and
b) the imprint of the positional changes of the animal's body over time in the animation ('pattern
image'; Figure 1B and Supplementary Video 1). These two pieces of data are then analyzed by
the Categorizer that comprises three submodules: the 'Animation Analyzer', the 'Pattern
Recognizer', and the 'Decision Maker' (Figure 1C). The Animation Analyzer analyzes all
spatiotemporal details in each animation, whereas the Pattern Recognizer evaluates the
animal's overall motion pattern in the paired pattern image (see details in "*Biologically plausible*
design of the Categorizer for holistic assessments"). The Decision Maker integrates the outputs
from the Animation Analyzer and Pattern Recognizer to determine the behavior category within
the specified time window.

Because it is based on deep neural networks, the Categorizer needs to be trained on
examples of ground truth (i.e., user-labeled data) for each behavior in order to conduct accurate
categorizations. To train the Categorizer, users only need to 'tell' the Categorizer what the
behaviors are by providing examples (animations and their paired pattern images generated by
LabGym; Figure 1B) of each behavior. To do so, users use *LabGym* to export unsorted example
files, and manually sort each pair of example files into folders created under the behavior name
(Figure 1D). Users then input these folders into *LabGym* to generate a labeled dataset for
training the Categorizer.

After the categorizations, the quantification module called the 'Quantifier' (Figure 1E)
uses information from the behavioral categories and the animal foregrounds to compute various
parameters for each behavior (e.g., duration, speed, etc.; Figure 1F). The output also includes

temporal raster plots of behavioral events (and their probabilities) and annotated videos for real-

time visualization of behavioral categorization.

Tracking multiple animals in diverse backgrounds for high-throughput and efficient

analyses

Behavioral experiments often involve behavior-irrelevant backgrounds—such as the enclosures

for holding the animals or changes in illumination during optogenetic manipulation—that renders

analyses inaccurate and inefficient. To address these issues, we designed a module in *LabGym*

that removes backgrounds but retains all the pixels representing the animals in each video

frame (i.e., foreground) (See also below in "*Biologically plausible design of the Categorizer for*

holistic assessments" for approaches for incorporating background information).

We reasoned that backgrounds in experimental conditions are often static, and that the

pixel value in each background location would be stable over time (Supplementary Figure 1A).

Therefore, we developed a method to find the stable value of each pixel in each video frame

and use these values to reconstruct the static background (Figure 2A and see **Methods**). The

static background is then subtracted from each frame to obtain the foreground, which

represents the animal (Figure 2B). Compared with two state-of-the-art background subtraction

methods, Local SVD Binary Pattern (LSBP) (Guo et al., 2016) and GSoC (Bradski, 2000)

(implemented in Google Summer of Code 2017), our stable-value detection method

reconstructed much cleaner backgrounds (with less traces of the animals) on behavioral videos

of multiple animals and in diverse experimental settings (Figure 2B and Supplementary Figure

1B). Notably, animals are reliably tracked by *LabGym* even in videos with unstable or shifting

illuminations (Supplementary Video 2) because the changes in background illumination (e.g.,

during optogenetic stimulation) can be reconstructed separately.

The tracking of each animal over time is performed based on the assumption that the

Euclidean distance between the centers of the same animal in consecutive frames are always

the smallest among all Euclidean distances between each possible pair of animal centers. Each

tracked animal is assigned by a unique identity (ID) that links to a matrix for storing all

information about that animal over time.

Multiple animals in the same enclosure sometimes collide with each other, which can

cause tracking errors. To make the analysis fully automatic, the current version of *LabGym*

excludes animals from tracking if users choose not to allow animal body entanglements (Figure

2C, Supplementary Videos 3 and **Methods**). This ensures that the analysis is only performed on

reliably tracked individual animals. If body entanglements are allowed, *LabGym* can also

analyze social behaviors in which animals have body contact (Figure 2C, Supplementary Videos

4 and **Methods**). In this scenario, merged animals are analyzed as one.

Taken together, the effective background subtraction and reliable animal tracking in

LabGym make it applicable to a wide range of behavioral videos without the need for

specialized recording equipment.

Biologically plausible design of the Categorizer for holistic assessments

We built the Categorizer with three submodules: Animation Analyzer, Pattern Recognizer, and

Decision Maker (Figure 3 and **Methods**), each of which consists of neural networks that suit the

design goals. Importantly, the two subnetworks—the Animation Analyzer and the Pattern

Recognizer—were designed to analyze two types of behavioral information that complement

each other, which ensures accurate categorizations of behavior through holistic assessment.

Animal behavior contains changes in both spatial and temporal dimensions. To learn the

spatial details of the animal in each time point (frame) during a behavior, the Animation Analyzer

uses time-distributed convolutional layers to analyze raw pixel values that represent the animal

foreground in each frame of an animation (Figure 3A). To learn about how these frame-wise

spatial details are organized along the temporal axis, the Animation Analyzer then uses

recurrent layers to compute temporal connectivity among the frame-wise spatial details

analyzed by time-distributed convolutional layers.

When an animal performs a behavior, the positional changes of its body and body parts

over time can form a pattern that is unique to the behavior. This pattern is key to behavioral

categorizations. To enable the Categorizer to capture distinct patterns, a "pattern image" was

generated for each animation to depict the animal's overall movement and the Pattern

Recognizer (that comprises convolutional layers) was designed to analyze the pattern image

(Figure 3B). In each pattern image, the contours of both the whole body and the body parts of

the animal in each frame of the animation were superimposed with gradually changed colors to

indicate their temporal sequences. This shows the animal's overall movement pattern during the

animation in a single 2-dimensional image to allow for efficient analyses.

Finally, the Decision Maker uses concatenating layers to integrate the outputs from both

the Animation Analyzer and the Pattern Recognizer, and passes the integrated information to

fully-connected (dense) layers for discriminating behavioral categories (Figure 3C).

Users can choose to include background in the animations as this sometimes can be

useful for behavioral categorization. For example, bedding in a mouse's mouth can indicate nest

building and, thus, including the bedding in the animations can facilitate the categorization of

nest building behavior (Figure 4A). Users can also choose whether to show the animals' body parts in the pattern images. Whole-body motion patterns are sometimes sufficient for distinguishing behaviors like curling versus uncoiling in *Drosophila* larvae, wing extension versus abdomen bending in adult flies, or walking versus rearing in rats (Figure 4B), while movement patterns of individual body parts are key for differentiating similar behaviors like facial grooming versus sniffing, or ear grooming versus sitting still in mice (Figure 4C).

The Categorizer accurately categorizes user-defined behaviors across species

Since the behavior types for the Categorizer to recognize are defined by users, rather than pre-defined in the program, the Categorizer can identify behaviors without restrictions on behavioral categories or animal species. To demonstrate this, we used *LabGym* to study relatively simple behaviors in *Drosophila* larvae and behaviors of increasing complexity in rodents. We first used *LabGym* to generate and label separate behavioral datasets for *Drosophila* larvae (Supplementary Video 5), rats (Supplementary Video 6), and mice (Supplementary Video 7). These datasets are publicly available (see **Methods**), can be used by others, and can be modified and expanded on by other users to suit their studies. The larva dataset contains the fewest categories with sufficient examples to cover almost all the variations of each behavior; the rat dataset contains a larger number of categories and more complex behaviors that encompass subtle distinctions between categories; the mouse dataset contains additional categories of even more complex behaviors. To test the robustness of *LabGym*, we also varied the amount of training examples used to categorize a given behavior, ranging from ~100 to ~800 pairs of animations and pattern images.

The Categorizer is customizable with different levels of complexity to address behaviors of various complexity. To provide a reference for customizing the Categorizer, which is a trial-and-error process, we trained several Categorizers with different complexity for each species, and determined the most suitable ones based on their overall accuracy of behavioral categorizations in training (Supplementary Table 1 and **Methods**). The selected Categorizer for each dataset was named as *LarvaNociceptor_Topview_30fps* (*LarvaN*), *RatAddiction_Topview_30fps* (*RatA*) and *MouseHomecage_Sideview_30fps* (*MouseH*). To test the accuracy and generalizability of these three Categorizers, we assessed their precisions, recalls (sensitivity), and F1 score (weighted precision and sensitivity, see **Methods**) for behavioral categorizations of videos that were not used for generating the training or validation dataset.

Complex deep neural networks might not be suitable for simple larval behaviors since
they would learn too many irrelevant details in simple datasets, resulting in reduced accuracy
(i.e., overfitting). In fact, *LarvaN* with simple complexity (Supplementary Table 1) achieved 97%
accuracy in frame-wise categorizations of 6 different behaviors elicited by noxious stimuli
(Figure 5A and Supplementary Video 8).
Rats show complex behaviors and the differences among them can be extremely subtle
due to the recording or experimental conditions (Supplementary Video 6). Despite these
challenges, the *RatA* performed reliably well in frame-wise differentiation of visually similar
behaviors such as body grooming versus face grooming, or head swaying versus in place
(Figure 5B and Supplementary Video 9).
Animals can develop new behaviors when experimental conditions change. For example,
mice in a cage equipped with a wheel tend to run on the wheel. The movement of running on
the wheel has features and kinematics that are distinct from running across a flat surface. Thus,
wheel-running might not be correctly identified by tools using predefined, generalized definition
of running. Additionally, a camera placed at the side of a cage will result in at least two
viewpoints of the mouse at it moves in the cage, a side view and a front view, which can greatly
complicate the automated analysis of behavior. Despite these challenges, *MouseH* achieved
excellent accuracy in frame-wise categorization of 20 behavioral categories (especially for
wheel-running, which achieved 100% precision and recall) with as few as 79 training examples
for some categories (Figure 5C and Supplementary Video 10). Although both were used for
categorizing rodent behaviors, *MouseH* trained on pattern images including the body parts
(Supplementary Videos 7) performed better than *RatA* trained on those excluding body parts
(Supplementary Videos 6; Figure 5B and 5C). This indicates that including the body parts in
pattern images can boost the learning efficiency of the Categorizer (Figure 4 and **Methods**).
Due to the data-driven nature of deep learning, the accuracy and generalizability of the
Categorizer largely depends on the amount, diversity, and labeling accuracy of the training
examples. Therefore, the performance of the Categorizers can be continuously improved over
time with the accumulation of well-labeled training examples.
29
The Quantifier provides quantitative measurements that are sufficient for capturing
subtle changes in diverse aspects of each behavior
To quantitatively measure the behavior intensity and the body kinematics during a behavior, we
designed a quantification module in *LabGym* called 'Quantifier'. The Quantifier calculates 14
parameters to quantify the behavior: count, latency, duration, angle, speed (distance traveled

per time unit), velocity (displacement per time unit), acceleration/velocity reduction, distance,

intensity (area), intensity (length), magnitude (area), magnitude (length), vigor (area) and vigor

(length) (Figure 6 and **Methods**). Additional parameters can be added in future versions of

LabGym. The definitions of these parameters are:

• The **count** is the summary of the behavioral frequencies, which is the occurrence

number of a behavior within the entire duration of analysis. Consecutive single

occurrences (at a single frame) of the same behavior are considered as one count.

• The **latency** is the summary of how soon a behavior starts, which is the time starting

from the beginning of the analysis to the time point that the behavior occurs for the first

time.

• The **duration** is the summary of how persistent a behavior is, which is the total time of a

behavior within the entire duration of analysis.

• The **angle** is the movement direction (against to the animal body axis) of the animal

during a behavior episode, which is the mean of all the included angle (θ) between

animal body axis and the movement direction during the time window (t_w) for

categorizing the behavior.

• The **speed** is the summary of how fast the animal moves when performing a behavior,

which is the total distance traveled (d) (between the two centers of mass of the animal)

during the time window (t_w) for categorizing the behavior divided by t_w.

• The **velocity** is the summary of how efficient the animal's movement is when performing

a behavior, which is the maximum displacement (dt) (between the two centers of mass

of the animal) divided by the time (t) that such displacement takes place.

• The **acceleration** / **velocity reduction** is the summary of how fast the animal's velocity

changes while performing a behavior, which is the difference between maximum velocity

(v_{max}) and minimum velocity (v_{min}) divided by the time (t_v) that such velocity change takes

place.

• The **distance** is the total distance traveled of the animal by performing a behavior within

the entire duration of analysis.

• The **intensity (area)** / **intensity (length)** is the summary of how intense a behavior is,

which is the accumulated proportional changes of the animal body area (a) / length (l)

between frames divided by the time window for categorizing the behaviors (t_w) when

performing a behavior.

- The **magnitude (area)** / **magnitude (length)** is the summary of the motion magnitude, which is the maximum proportional change in animal body area (a) or length (l) when performing a behavior.
- The **vigor (area)** / **vigor (length)** is the summary of how vigorous a behavior is, which is the magnitude (area) / magnitude (length) divided by the time (t_a or t_l) that such a change takes place.

To test how well the Quantifier performs in quantifying behaviors in diverse animal species, we applied *LabGym* to experiments in larvae, rats, and mice, in which the quantitative measurements of behaviors was difficult to achieve with previous methods.

Drosophila larva behavior:

Previous studies show that optogenetic inhibition of larval leucokinin (LK) neurons slightly impaired larval behaviors elicited by nociceptor activation (Hu et al., 2020). In these experiments, both the subtle changes in behaviors and the shifting illumination caused by optogenetic manipulations posed challenges for automatic behavioral analysis. We thus tested whether *LabGym* was able to capture the behavioral changes in these experiments (Supplementary Video 11). Compared with the control group, larvae in the LK inhibition group showed a modest reduction in the probability of body rolling behavior, which is a characteristic nociceptive response (Hwang et al., 2007; Tracey et al., 2003) (Figure 7A and 7B). No change was observed in the latency of the response onset after optogenetic stimulation of nociceptors (Figure 7C). Moreover, LK inhibition caused a slight yet significant reduction in rolling velocity (Figure 7D) and duration (Figure 7E). We also compared several quantitative measurements of different behaviors and found that the results indeed reflect the ethological meaning of these behaviors. For example, rolling achieves higher speed than crawling (Figure 7F), supporting the notion that rolling was a more efficient way than crawling to escape from harm (Hwang *et al.*, 2007; Tracey *et al.*, 2003). Similarly, despite their spatial similarity, curling was more vigorous than turning (Figure 7G), echoing the fact that larvae typically perform curling under noxious stimuli but tend to turn in in response to milder situation like gentle touch or exploration (Hwang *et al.*, 2007; Jovanic et al., 2016; Lahiri et al., 2011; Tracey *et al.*, 2003). Thus, *LabGym* is sufficiently sensitive to detect modest effects of in vivo neuronal inhibition on behavior, and it does so on videos with shifting illumination.

When *Drosophila* larvae are exposed to sound, they display a startle response that is characterized by the cessation of crawling ("freezing"), retraction of their heads ("hunching"),

and excessive turning movements (Zhang et al., 2013). To test whether *LabGym* could identify

the subtle differences in behaviors elicited by different intensities of sound, we trained another

Categorizer (*LarvaMechanosensor_TopView_30fps*) with examples of sound-elicited behaviors

in *Drosophila* larvae (Supplementary Videos 5) and used it for analysis (Supplementary Video

12). This Categorizer has the same complexity as *LarvaN*. Compared to larvae exposed to 80

decibel (dB) white noise, those exposed to 94 dB showed a similar hunching probability (Figure

7H and 7I), but the magnitude of head retraction during hunching was enhanced (Figure 7I).

Furthermore, larvae exposed to 94 dB showed more frequent turning (Figure 7K) and prolonged

freezing (i.e., immobility; Figure 7I), which resulted in significant reductions in crawling distance

(Figure 7M). However, when larvae froze, the reduction in the velocity of their movement was

similar between the two groups (Figure 7N). Thus, *LabGym* was able to capture quantitative

differences in the behaviors induced by auditory stimuli of different intensities.

Rodent behavior:

It is well-established that drugs like amphetamine, produce dose-dependent increases in

psychomotor activity, and that the intensity of these behaviors increases (i.e., sensitizes) with

repeated drug exposure (Ellinwood and Balster, 1974; Ferrario *et al.*, 2005). Psychomotor

activity in rats encompasses a wide range of behaviors that include rearing, hyper-locomotion,

and stereotyped, repetitive head movements (i.e., stereotypy). Psychomotor activity is most

often quantified using non-parametric rating scales based on clusters of behavioral features and

their intensities (Ellinwood and Balster, 1974; Ferrario *et al.*, 2005). To date, no automated

systems have been able to sufficiently categorize or quantify psychomotor activity, despite early

efforts that predate machine learning (Flagel and Robinson, 2007).

LabGym automatically captured dose-dependent changes in key behavioral features of

psychomotor activity (Figure 8A and Supplementary Video 13). These include rearing (Figure

8B), locomotion (Figure 8C), and head movements (persistent head movements indicating

stereotypy; Figure 8D), following systemic amphetamine administration. Impressively, *LabGym*

even captured stereotypy interspersed with small bursts of locomotion—an indication of a strong

psychomotor response at high dose of amphetamine (e.g., Figure 8A and 8C, 5.6 mg/kg dose

group in amphetamine baseline group). *LabGym* also captured quantitative changes in these

behaviors across the development of sensitization, in which repeated injection of amphetamine

resulted in more intense behavioral responses to the same dose of drug (Figure 8E-8H). In

contrast, the intensity of spontaneously occurring behaviors such as body/face grooming were

stable across time in rats treated with repeated saline (Figure 8I-8K).

When an animal ages, its ethogram might not change, yet the quantitative

measurements of its behaviors (e.g., running speed) may decline. The speed of a mouse

running on a wheel is typically measured by tracking the rotation of the wheel itself, which poses

a challenge for computational tools designed to only track animals. Nevertheless, *LabGym* not

only categorized the all the behaviors of interest for both younger (32-week) and older mice (80-

week) in a cage with a running wheel (Figure 8L and Supplementary Video 14), but also

captured the decline in movement speed of the older versus the young mice by a parameter

(vigor_area) that summarizes the areal changes per unit time of the animal's body (Figure 8M, 6

and **Methods**).

Taken together, these results demonstrate that the Quantifier reliably and precisely

captures quantitative changes in behaviors across species, and in a wide range of settings and

experimental conditions.

GUI with optional settings for user-friendly usage

To make *LabGym* adaptable to various experimental settings and accessible to users without

requiring a computer programming background, we implemented a GUI. This includes 4

functional units: Generate Datasets, Train Networks, Test Networks and Analyze Behaviors.

The Generate Datasets unit (Supplementary Figure 2) is for batch processing of videos

to generate visualizable behavioral data (animations and paired pattern images) that can be

used to establish benchmark datasets for training the neural networks. The Train Networks unit

(Supplementary Figure 3) is for preparing training datasets and training Categorizers at various

levels of complexity. The Test Networks unit (Supplementary Figure 4) is for testing the

accuracy and generalizability of a trained Categorizer in an unbiased manner. The Analyze

Behaviors unit (Supplementary Figure 5) is for batch-processing of videos for behavioral

analysis. It generates an annotated video (Supplementary Videos 8-14), a raster plot of

behavioral events (displayed in user-defined colors) and their occurring probabilities (by color

intensities) for each animal in one batch of analysis (Figure 7A, 7H, 8A and 8L), and a

spreadsheet in Excel or CSV format containing individual behavioral measures for each

behavior of each animal.

Notably, the flexible options in *LabGym*, especially the customizable neural network

complexity, allows it to achieve fast processing speed in training networks and in analyzing

behaviors on commonly used laptops that do not have GPUs (Supplementary Table 2). This is

difficult to achieve for tools using fixed and complex neural networks (Bohnslav *et al.*, 2021;

Mathis *et al.*, 2018).

Benchmark comparison with the existing standards

The combination of deep learning-based tracking tools and the post-hoc behavior analysis tools

can achieve multi-animal tracking, classification and quantification of user-defined behaviors in

different animal species, while *LabGym* can achieve the same goal in one tool. We directly

benchmarked the performance of *LabGym* against that of the combination of DeepLabCut (DLC)

(Mathis *et al.*, 2018) and B-SOiD (Hsu and Yttri, 2021) ("DLC+B-SOiD"), which represent the

existing standards and also have GUIs for accessibility to users without requiring computer

programming knowledge.

All the three tools can be iteratively optimized by adding more training data and refining

the data labeling. Therefore, the goal of the benchmark comparison is not to explore how to

modify the training datasets to achieve the best performance of these tools, but to compare their

performance with the same training resources and labor workload (using a fixed 30-minute for

manual labeling and sorting for both *LabGym* and DLC+B-SOiD). To achieve this, we used

videos in which animals display behaviors of interest that differ in intensity. We then determined

whether each approach (DLC+B-SOiD and *LabGym*) can detect the behaviors of interest, and

how results compare across these approaches (Figure 9A). Specifically, the benchmarking

training dataset used contained 4 videos (2-minute duration at 30 fps) covering major types of

rat behavior in psychomotor activity experiments: locomotion, rearing, repeated head

movements and other background behaviors such as resting (in place) and orientating. The

labor for DLC+B-SOiD was spent on labeling animal body parts in extracted frames for tracking

(~25 minutes) and assigning behavior names for behavior classification (~5 minutes), while that

in *LabGym* was spent on sorting behavior data for behavior classification (~30 minutes). The

testing dataset contained 3 videos (1-minute duration at 30 fps per video):

1) A video (referred as 'Base') that was randomly selected and trimmed from

'Amphetamine (baseline)' in Figure 8A, in which the rats were exposed to 5.6 mg/kg

amphetamine for the first time. The rat in this video showed psychomotor activities such as

frequent rearing and locomotion.

2) A video (referred as 'Post') that was randomly selected and trimmed from

'Amphetamine (post treatment)' in Figure 8A, in which the rats were exposed to 5.6 mg/kg

amphetamine after experiencing repeated amphetamine exposure (i.e., a sensitizing regimen).

The rat in this video showed intensified psychomotor activity such as frequent stereotypy

(repeated head moving).

3) A video from (Ferrario *et al.*, 2005) in which testing occurred in a different enclosure

than that used in the training videos. This video was used to test the generalizability of these

tools.

Comparisons were performed on two key aspects: the total time cost and the end

accuracy of identifying and quantifying behaviors of interest. We first input the 4 training videos

(8 minutes in total at 30 fps) into DLC to extract frames (~2 minutes) and labeled 6 body parts in

each frame (~25 minutes). We then used these labeled frames to train a DLC model (~8 hours).

This resulted in only 2.83 pixels of error in evaluation, indicating a successful training

(Supplementary Figure 6A). Next, we ran the trained DLC model on the 4 training videos to get

the frame-wise positions of the 6 body parts (~15 minutes). These outputs were then fed into B-

SOiD to train a B-SOiD model for behavior classification (~10 minutes). B-SOiD uses

unsupervised approach to form behavior groups and outputs video examples for each group,

which is a convenient way for users to associate these computer-decided groups with

ethologically meaningful behaviors (~5 minutes). We tested different number of behavior groups

from ~100 to 5 groups and inspected the video examples for each group. We found that the 5-

group were the best fit since: (1) others resulted in either too many different groups that turned

out to be the same behavior that was expressed in different ways, or a single group containing

different behaviors that could not be assigned to a single ethologically meaningful behavior type;

and (2) the 5-group B-SOiD model showed 5 well-separated clusters (Supplementary Figure

6B). We then labeled each group as a single behavior type based on the most frequently

observed behavior among the videos in that group (Supplementary Figure 6C). Finally, we ran

the trained DLC model on the 3 testing videos (3 minutes in total at 30 fps) to produce the

frame-wise body part position output and fed that into the trained B-SOiD model for behavior

classification and quantification (~5 minutes).

LabGym does not require labeling or training to track the animals; users only need to

specify a time window in a video for background extraction. Thus, as described above, we input

the 4 training videos into *LabGym* to generate data pairs containing animations of user-defined

duration and their paired pattern images (~5 minutes). We then sorted these data pairs into

different folders according to user-defined behaviors of interest (~30 minutes). Next, we input

these folders into *LabGym* to train a Categorizer for behavior classification (~10 minutes).

Finally, we ran *LabGym* using the trained Categorizer on the 3 testing videos for behavior

classification and quantification (~5 minutes).

The total time to set up and run DLC+B-SOiD on 8-minute training videos and 3-minute

testing videos at 30 fps was about 9 hours (including 0.5 hour of labor), while that for *LabGym*

was less than 1 hour (including 0.5 hour of labor; Figure 9B). To establish the 'ground truth' for

determining the end accuracy of frame-wise identification of behaviors in the 3 testing videos,

we manually annotated the behavior types in each frame of the 3 testing videos (1,800 frames

per video). We then compared the frame-wise behavior classification outputs of B-SOiD or

LabGym with this ground truth. We found that both B-SOiD and *LabGym* accurately identified

rearing and locomotion in the 'Base' video ('Amphetamine (baseline)'; Figure 9C and

Supplementary Video 15). However, B-SOiD failed to identify head movements in 'Post' video

('Amphetamine (post treatment)'), whereas *LabGym* achieved good accuracy in identifying this

behavior (Figure 9D and Supplementary Video 16).

Both B-SOiD and *LabGym* provided quantitative measurements of the animal body

kinematics during locomotion (Figures 9E and 9F), but only *LabGym* captured the real

quantitative differences in behavior between rats given a single vs. repeated amphetamine

exposure (Figure 9F). Specifically, after repeated exposure the frequency of locomotion was

dramatically decreased while repeated head movements (i.e., in-placed stereotyped head

movements) were increased (Supplementary Videos 15 and 16), causing a reduction in

locomotion distance. This is a classic pattern that typifies the expression of psychomotor

sensitization (Carr et al., 2020; Ferrario *et al.*, 2005). Additionally, it is noteworthy that the speed

of locomotion was similar across groups. This highlights the need to quantify a behavior to

accurately measure effects of psychostimulant drugs on behavior (see (Ferrario *et al.*, 2005) for

additional discussion).

When the testing was applied to a video made in a different enclosure, the Categorizer

in *LabGym*, which was trained on only 49~119 training data pairs for each behavior type, was

able to generalize the behavioral diction with only a modest drop in accuracy (Figure 9G and

Supplementary Video 17). In contrast, B-SOiD was not able to generalize well (Figure 9G).

It is possible that B-SOiD may be able to accurately identify the "head movement"

behavior in the testing videos if additional training data were provided. Alternatively, it is

possible that the pre-defined features (body length, angle, and speed) used by B-SOiD to form

behavioral groups might not be useful for distinguishing head movements from other behaviors.

Nevertheless, the purpose here was to directly compare approaches using the same training

resource. Both approaches were able to accurately track rearing, but only *LabGym* was able to

generalize to another testing enclosure, and capture behaviors specific to psychostimulant drug

effects.

DISCUSSION

LabGym is a unique computational tool that enables a generalizable solution to automated behavioral analysis. The flexibility and trainability of this tool allows it to be applied to very broad variety of species, behaviors and questions, and to be iteratively refined towards a particular set of conditions over time. This tool opens opportunities for automatic, high-throughput, targeted and customizable analyses of behaviors without the cost of commercial products, which will greatly facilitate the process of mechanistic discoveries in biological research. The standardized, visualizable behavioral datasets generated by *LabGym* will also benefit research in neuroscience, behavioral science, and computer science by helping to standardize procedures within and across research groups, and by providing long-term references for future studies to build on.

Innovations in *LabGym*

Firstly, we developed a new method for effective background subtraction in *LabGym* that outperforms the state-of-the-art background subtraction approaches. This new method enables accurate tracking of animal bodies without the need of manual labeling and model training, which significantly saves the human labor and accelerates the analysis.

Secondly, unlike other tools using fixed and complex neural networks (Bohnslav *et al.*, 2021; Mathis *et al.*, 2018) that are slow to train and hard to generalize, we designed a new neural network structure in *LabGym* with customizable combinations of Animation Analyzer, Pattern Recognizer and Decision Maker. This allows users to customize the behavior categorization to suit behavior data with different complexity and enables fast running speed in even common laptops without GPUs. This new design also ensures the effectiveness in holistic assessment on both the complete spatiotemporal details in a behavior and the overall movement pattern that is unique to the behavior. The holistic assessment on behavioral information in *LabGym* does not rely on pre-defined features and does not require intermediate steps of extracting features, which minimizes the information loss.

Thirdly, we developed a new method for evaluating the overall pattern of the animal's movement (including both the whole body and limbs) during a behavior. We designed the "pattern image" to exhibit the animal's motion pattern of both the whole body and body parts efficiently and precisely. We then designed the Pattern Recognizer in Categorizer to analyze the pattern image. This method is more efficient and precise than current approaches for assessing

motion flows that operate on every pixel in video frames to generate optical flows (Bohnslav *et*

al., 2021).

Lastly, *LabGym* is "end-to-end": it provides a GUI and does not need intermediate or

post-hoc steps that require users to have expertise in computation or programming. Instead,

users only need to recognize and sort the animations and their paired pattern images as

examples of a given behavior to 'teach' *LabGym* to identify the behavior. The behavior is then

automatically quantified by *LabGym*.

LabGym provides a way for users to generate standardized, visualizable behavioral

datasets

In behavioral categorization based on deep learning approaches, the accuracy depends on the

quality of training datasets, which should be diverse and well labeled (LeCun *et al.*, 2015; Roh

et al., 2021; Schmidhuber, 2015). In the computer vision field, high-quality datasets, such as

ImageNet (Deng et al., 2009), have been established to facilitate the training of deep neural

networks and to benchmark different algorithms. In neuroscience, however, such benchmark

datasets for behaviors in model organisms are largely absent. A source of this problem is the

lack of tools that can help researchers to generate standardized, visualizable behavioral

datasets that are easy to label and to share. Although transfer learning that applies acquired

network weights to new categorization problems can potentially be used to compensate for the

lack of species-specific behavioral datasets, its efficiency relies on the similarity between

existing datasets used for pretraining the networks and the new data (Yosinski et al., 2014).

Current tools using transfer learning for behavioral categorizations (Bohnslav *et al.*, 2021)

utilized network weights that were acquired from human behavioral datasets such as Kinetics-

700 (Carreira et al., 2019). However, human behaviors are intrinsically different from the

behaviors of model organisms, especially the soft-bodied ones. Therefore, species-specific

behavioral datasets are still in critical demand for accurate behavioral categorizations in model

organisms.

LabGym addresses the lack of such benchmark datasets by providing a way for users to

generate standardized, visualizable behavioral datasets that to compare their categorization

(labeling) of behavior and refine the labeling. Importantly, such expert-labeled behavioral

datasets, if shared in the research community, not only ensure the accurate and reproducible

executions of categorization criteria across different experimenters or trials, but also provide

benchmarks for comparing different algorithms, and references for future studies to build on.

Limitations of *LabGym*

LabGym does not require labeling or training to track the animals. In turn, it does have preferred settings for video recording. Although *LabGym* does not have restriction on the kind of background/arena/camera to use, the background/arena/camera needs to be static during the period for behavior analysis. The illumination can have dark-to-bright or bright to dark transitions but need to be stable before and after the transitions during the period for behavior analysis. Animals are expected to present some locational changes during video recording, instead of being immobile all the time. The time window during which animals are moving can be used for background extraction in *LabGym*. This requirement in video recording can be alleviated by implementing mask region-based convolutional neural networks (mask R-CNN) (He et al., 2017) in future versions of *LabGym*. Mask R-CNN can be used to segment animals from dynamic backgrounds (i.e., the backgrounds continually change over time). Currently, when analyzing animals in dynamic backgrounds, users may first consider using the mask R-CNN (https://github.com/matterport/Mask_RCNN) (Abdulla, 2017) to obtain animal-masked videos, and then use the current version of *LabGym* to analyze such videos. Additional labeling and training efforts will be needed in such cases.

The current version of *LabGym* is not optimized for analyzing social behaviors. In social behaviors, animals have body contacts. The animal identity reassignment after animals re-separate might not be accurate in *LabGym*. The accuracy of animal identity reassignment can be improved by implementing mask R-CNN in future versions of *LabGym*.

ACKNOWLEDGMENTS

We thank Drs. Daniel Leventhal, Hanchuan Peng, and Guoqiang Yu for helpful comments on

earlier versions of the manuscript. This research was supported by National Institutes of Health

(NIH) to B.Y. (R01NS104299 and R01EB028159), and C.R.F. (R01-DK106188, R01-DK115526,

R01-DA044204). *Drosophila* Stocks from the Bloomington *Drosophila* Stock Center (NIH

P40OD018537) were used in this study. The content is solely the responsibility of the authors

and does not necessarily represent the official views of the NIH.

AUTHOR CONTRIBUTIONS

Y.H. and B.Y. conceived the project. Y.H. designed and programmed *LabGym* with the

guidance of J.Z. and B.Y.. Y.H., K.I., H.W., C.R.F., A.G., and B.W. performed the behavioral

experiments in *Drosophila* larvae (Y.H, K.I. and H.W.), rats (C.R.F.), and mice (A.G. and B.W.).

Y.H., K.I., H.W., and Y.X. generated and labelled the dataset for larval behaviors (Y.H., K.I.,

H.W. and Y.X.), rat behaviors (Y.H.), and mouse behaviors (Y.H. and A.G.). Y.H. performed the

computational analyses and the validation of behavioral categorizations with the assistance

from C.R.F., A.G., B.W., K.I., and H.W.. Y.H, A.B.M. and R.B.I. performed the benchmark

comparison between *LabGym* and DeepLabCut+B-SOiD. Y.H., C.R.F., B.W., J.Z., and B.Y.

wrote the manuscript.

DECLARATION OF INTERESTS

The authors declare no competing interests.

REFERENCES

Abdulla, W. (2017). Mask R-CNN for object detection and instance segmentation on Keras and TensorFlow. GitHub repository.

Altimus, C.M., Marlin, B.J., Charalambakis, N.E., Colon-Rodriquez, A., Glover, E.J., Izbicki, P., Johnson, A., Lourenco, M.V., Makinson, R.A., McQuail, J., et al. (2020). The Next 50 Years of Neuroscience. J Neurosci *40*, 101-106. 10.1523/Jneurosci.0744-19.2019.

Berman, G.J., Choi, D.M., Bialek, W., and Shaevitz, J.W. (2014). Mapping the stereotyped behaviour of freely moving fruit flies. J R Soc Interface *11*. 10.1098/rsif.2014.0672.

Bohnslav, J.P., Wimalasena, N.K., Clausing, K.J., Dai, Y.Y., Yarmolinsky, D.A., Cruz, T., Kashlan, A.D., Chiappe, M.E., Orefice, L.L., Woolf, C.J., and Harvey, C.D. (2021). DeepEthogram, a machine learning pipeline for supervised behavior classification from raw pixels. Elife *10*. 10.7554/eLife.63377.

Bradski, G. (2000). The OpenCV library. Dr Dobbs J *25*, 120-+.

Carr, C.C., Ferrario, C.R., and Robinson, T.E. (2020). Intermittent access cocaine self-administration produces psychomotor sensitization: effects of withdrawal, sex and cross-sensitization. Psychopharmacology (Berl) *237*, 1795-1812. 10.1007/s00213-020-05500-4.

Carreira, J., Noland, E., Hillier, C., and Zisserman, A. (2019). A short note on the kinetics-700 human action dataset. arXiv preprint arXiv:1907.06987.

Datta, S.R., Anderson, D.J., Branson, K., Perona, P., and Leifer, A. (2019). Computational Neuroethology: A Call to Action. Neuron *104*, 11-24. 10.1016/j.neuron.2019.09.038.

Deng, J., Dong, W., Socher, R., Li, L.J., Li, K., and Li, F.F. (2009). ImageNet: A Large-Scale Hierarchical Image Database. Cvpr: 2009 Ieee Conference on Computer Vision and Pattern Recognition, Vols 1-4, 248-255. DOI 10.1109/cvpr.2009.5206848.

Denisov, G., Ohyama, T., Jovanic, T., and Zlatic, M. (2013). Model-based Detection and Analysis of Animal Behaviors using Signals Extracted by Automated Tracking. pp. 175-181.

Egnor, S.E., and Branson, K. (2016). Computational Analysis of Behavior. Annu Rev Neurosci *39*, 217-236. 10.1146/annurev-neuro-070815-013845.

Ellinwood, E.H., Jr., and Balster, R.L. (1974). Rating the behavioral effects of amphetamine. Eur J Pharmacol *28*, 35-41. 10.1016/0014-2999(74)90109-5.

Ferrario, C.R., Gorny, G., Crombag, H.S., Li, Y.L., Kolb, B., and Robinson, T.E. (2005). Neural and behavioral plasticity associated with the transition from controlled to escalated cocaine use. Biol Psychiat *58*, 751-759. 10.1016/j.biopsych.2005.04.046.

Flagel, S.B., and Robinson, T.E. (2007). Quantifying the psychomotor activating effects of cocaine in the rat. Behav Pharmacol *18*, 297-302. 10.1097/FBP.0b013e3281f522a4.

Fukushima, K. (1980). Neocognitron - a Self-Organizing Neural Network Model for a Mechanism

of Pattern-Recognition Unaffected by Shift in Position. Biol Cybern *36*, 193-202. Doi

10.1007/Bf00344251.

Guo, L.L., Xu, D., and Qiang, Z.P. (2016). Background Subtraction using Local SVD Binary

Pattern. Ieee Comput Soc Conf, 1159-1167. 10.1109/Cvprw.2016.148.

He, K., Gkioxari, G., Dollár, P., and Girshick, R. (2017). Mask r-cnn. pp. 2961-2969.

Hsu, A.I., and Yttri, E.A. (2021). B-SOiD, an open-source unsupervised algorithm for

identification and fast prediction of behaviors. Nature Communications *12*, 5188.

10.1038/s41467-021-25420-x.

Hu, Y.J., Wang, C.C., Yang, L.M., Pan, G., Liu, H., Yu, G.Q., and Ye, B. (2020). A Neural Basis

for Categorizing Sensory Stimuli to Enhance Decision Accuracy. Curr Biol *30*, 4896-+.

10.1016/j.cub.2020.09.045.

Hwang, R.Y., Zhong, L.X., Xu, Y.F., Johnson, T., Zhang, F., Deisseroth, K., and Tracey, W.D.

(2007). Nociceptive neurons protect Drosophila larvae from parasitoid wasps. Curr Biol *17*,

2105-2116. 10.1016/j.cub.2007.11.029.

Jovanic, T., Schneider-Mizell, C.M., Shao, M., Masson, J.B., Denisov, G., Fetter, R.D., Mensh,

B.D., Truman, J.W., Cardona, A., and Zlatic, M. (2016). Competitive Disinhibition Mediates

Behavioral Choice and Sequences in Drosophila. Cell *167*, 858-+. 10.1016/j.cell.2016.09.009.

Kabra, M., Robie, A.A., Rivera-Alba, M., Branson, S., and Branson, K. (2013). JAABA:

interactive machine learning for automatic annotation of animal behavior. Nat Methods *10*, 64-

U87. 10.1038/Nmeth.2281.

Klibaite, U., Berman, G.J., Cande, J., Stern, D.L., and Shaevitz, J.W. (2017). An unsupervised

method for quantifying the behavior of paired animals. Phys Biol *14*. ARTN 015006

10.1088/1478-3975/aa5c50.

Klibaite, U., and Shaevitz, J.W. (2020). Paired fruit flies synchronize behavior: Uncovering social

interactions in Drosophila melanogaster. PLoS Comput Biol *16*, e1008230.

10.1371/journal.pcbi.1008230.

Lahiri, S., Shen, K.L., Klein, M., Tang, A.J., Kane, E., Gershow, M., Garrity, P., and Samuel,

A.D.T. (2011). Two Alternating Motor Programs Drive Navigation in Drosophila Larva. Plos One

6. ARTN e23180

10.1371/journal.pone.0023180.

LeCun, Y., and Bengio, Y. (1995). Convolutional networks for images, speech, and time series.

The handbook of brain theory and neural networks *3361*, 1995.

LeCun, Y., Bengio, Y., and Hinton, G. (2015). Deep learning. Nature *521*, 436-444.

10.1038/nature14539.

Marks, M., Jin, Q., Sturman, O., von Ziegler, L., Kollmorgen, S., von der Behrens, W., Mante, V.,

Bohacek, J., and Yanik, M.F. (2022). Deep-learning-based identification, tracking, pose

estimation and behaviour classification of interacting primates and mice in complex

environments. Nature Machine Intelligence *4*, 331-340. 10.1038/s42256-022-00477-5.

Mathis, A., Mamidanna, P., Cury, K.M., Abe, T., Murthy, V.N., Mathis, M.W., and Bethge, M.

(2018). DeepLabCut: markerless pose estimation of user-defined body parts with deep learning.

Nature Neuroscience *21*, 1281-1289. 10.1038/s41593-018-0209-y.

Nilsson, S.R., Goodwin, N.L., Choong, J.J., Hwang, S., Wright, H.R., Norville, Z.C., Tong, X.,

Lin, D., Bentzley, B.S., and Eshel, N. (2020). Simple Behavioral Analysis (SimBA)–an open

source toolkit for computer classification of complex social behaviors in experimental animals.

BioRxiv.

Pereira, T.D., Aldarondo, D.E., Willmore, L., Kislin, M., Wang, S.S.H., Murthy, M., and Shaevitz,

J.W. (2019). Fast animal pose estimation using deep neural networks. Nat Methods *16*, 117-125.

10.1038/s41592-018-0234-5.

Pereira, T.D., Shaevitz, J.W., and Murthy, M. (2020). Quantifying behavior to understand the

brain. Nat Neurosci *23*, 1537-1549. 10.1038/s41593-020-00734-z.

Roh, Y., Heo, G., and Whang, S.E. (2021). A Survey on Data Collection for Machine Learning:

A Big Data-AI Integration Perspective. Ieee T Knowl Data En *33*, 1328-1347.

10.1109/Tkde.2019.2946162.

Schmidhuber, J. (2015). Deep learning in neural networks: An overview. Neural Networks *61*,

85-117. 10.1016/j.neunet.2014.09.003.

Segalin, C., Williams, J., Karigo, T., Hui, M., Zelikowsky, M., Sun, J.J., Perona, P., Anderson,

D.J., and Kennedy, A. (2021). The Mouse Action Recognition System (MARS) software pipeline

for automated analysis of social behaviors in mice. Elife *10*. 10.7554/eLife.63720.

Tracey, W.D., Wilson, R.I., Laurent, G., and Benzer, S. (2003). painless, a Drosophila gene

essential for nociception. Cell *113*, 261-273. Doi 10.1016/S0092-8674(03)00272-1.

Wiltschko, A.B., Johnson, M.J., Iurilli, G., Peterson, R.E., Katon, J.M., Pashkovski, S.L., Abraira,

V.E., Adams, R.P., and Datta, S.R. (2015). Mapping Sub-Second Structure in Mouse Behavior.

Neuron *88*, 1121-1135. 10.1016/j.neuron.2015.11.031.

Yosinski, J., Clune, J., Bengio, Y., and Lipson, H. (2014). How transferable are features in deep

neural networks ? Adv Neur In *27*.

Zhang, W., Yan, Z.Q., Jan, L.Y., and Jan, Y.N. (2013). Sound response mediated by the TRP

channels NOMPC, NANCHUNG, and INACTIVE in chordotonal organs of Drosophila larvae. P

Natl Acad Sci USA *110*, 13612-13617. 10.1073/pnas.1312477110.

1
2 **Figure 1. The pipeline of *LabGym*.**

3 **A**. Static backgrounds are removed in each video frame and individual animals are tracked.

4 **B**. At each frame of the video, *LabGym* generates a pair of data for each animal, which

5 comprises a short animation spanning a user-defined time window and its paired pattern image

6 representing the movement pattern of the animal within the animation.

C. The Categorizer uses both the animations and pattern images to categorize behaviors during

the user-defined time window (i.e., the duration of the animation). The Categorizer comprises

three submodules, the 'Animation Analyzer', the 'Pattern Recognizer', and the 'Decision Maker'.

The Animation Analyzer analyzes all spatiotemporal details in the animation for each animal

whereas the Pattern Recognizer evaluates the overall movement patterns in the paired pattern

image. The Decision Maker integrates the outputs from both the Animation Analyzer and the

Pattern Recognizer to determine which user-defined behavioral category is present in the

animation for each animal.

D. The animations and pattern images can be exported to build visualizable, sharable

benchmark datasets for training the Categorizer.

E and **F**. After the behavioral categorizations, the quantification module ('Quantifier') uses

information from the behavioral categories and the animal foregrounds to compute specific

quantitative measurements for different aspects of each behavior.

Figure 2. *LabGym* removes backgrounds and accurately tracks multiple animals.

A. The stable value of each pixel is used to reconstruct the static background for a video.

B. Examples of video frames showing that animal foregrounds are clearly presented after

removing the static backgrounds that are constructed by our method of stable-value detection.

C. Examples of video frames showing that *LabGym* accurately track multiple animals performing

either non-interactive or social behavior. Left: non-interactive behavior, in which animal (larva)

contact is not allowed; right: social interaction in which animal (adult fly) contact is allowed.

Figure 3. Biological plausible design of the Categorizer.

A. The Animation Analyzer first uses time-distributed convolutional layers to compute all frame-wise spatial details of an animation, and then uses recurrent layers (long short-term memory, LSTM) to compute the temporal connectivity among the frame-wise spatial details.

B. The Pattern Recognizer uses convolutional layers to analyze the pattern image, which is

superimposed contours of animal body (and body parts) along the temporal sequence (indicated

by gradually changed color) during a behavior.

C. The Decision Maker uses concatenating layers to integrate the outputs from both the

Animation Analyzer and the Pattern Recognizer, and then passes the integrated information to

dense layers for concluding the behavioral categories.

Figure 4

A

Mouse nest building

B

Larva curling

Larva uncoiling

Fly abdomen bending

Fly wing extension

Rat walking

Rat rearing

C

Mouse facial grooming

Mouse sniffing

Mouse ear grooming

Mouse sitting

Figure 4. *LabGym* generates stand-alone, visualizable behavioral benchmark datasets.

A. Example frames of an animation (background is included) and its paired pattern image,

showing nest building behavior of a mouse.

B. Example frames of animations and their paired pattern images (body parts are not included),

showing (from top to bottom) larva curling and uncoiling, fly abdomen bending (copulation

attempt) and wing extension (courtship song), and rat walking and rearing.

C. Example frames of animations and their paired pattern images (body parts are included),

showing (from top to bottom) mouse facial grooming, sniffing, ear grooming, and sitting.

Figure 5

A LarvaNociceptor_Topview_30fps (overall accuracy: 0.97)

Training example amount (in pairs) (Tr N): 492 457 500 413 329 308

B RatAddiction_Topview_30fps (overall accuracy: 0.90)

Tr N: 369 229 551 551 539 552 561 512

C MouseHomecage_Topview_30fps (overall accuracy: 0.96)

Tr N: 177 90 145 123 200 373 148 89 368 500 317 331 424 116 232 308 79 217 392 240

Figure 5. Accurate frame-wise categorizations of various user-defined behaviors by

different Categorizers.

A-**C**. Bars showing the in-practice test metrics (overall accuracy, precisions, recalls and F1

scores) for *LarvaN* (**A**), *RatA* (**B**) and *MouseH* (**C**) in video analysis. 7 videos for larvae (10

larvae per video), 10 videos for rats (1 rat per video), and 2 videos for mice (1 mouse per video)

that were not used for generating the training datasets were used to generate the testing data

(animations and their paired pattern images). 2162 pairs (for larvae), 4828 pairs (rats), and

12653 pairs (mice) of data were randomly selected (data with visibly missing body parts were

excluded) and then were sorted by experimenters into different folders under user-defined

behavioral names (building ground truth testing datasets). The ground truth testing datasets

were then used to test the categorization metrics of the 3 Categorizers. In this way,

experimenters were blind to the predictions of the 3 Categorizers when sorting data to build the

ground truth testing datasets and thus the tests were unbiased.

2   **Figure 6. The Quantifier analyzes diverse aspects of a behavior.**

3   A schematic that shows the process of frame-wise categorizations on behaviors, and how the

4   Quantifier uses the information of behavioral categories and animal foregrounds to calculate 14

5   behavioral parameters in *LabGym*.

Figure 7

Figure 7. The Quantifier captures subtle changes in the behavior of *Drosophila* larvae.

A and **H**. Raster plots showing the frame-wise categorizations by: **A**. *LarvaN* on control (*w*; *LK-*GAL4 / +; *TrpA1*-QF, QUAS-ChR2[T159C], UAS-jRCaMP1b / +) and LK neuron inhibition (*w*; *LK-*

GAL4 / +; *TrpA1*-QF, QUAS-ChR2^{T159C}, UAS-GtACR1 / +) groups after nociceptors were

optogenetically activated (blue dashed line indicates the stimulation onset); **H**.

LarvaMechanosensor_Topview_30fps on larvae (*Caton S*) under different intensities of sound

stimuli (black dashed line indicates the stimulation onset). Color bars: behavioral categories

(color intensities indicate the behavioral probabilities). X-axis represents the time and Y-axis

represents the larval IDs.

B-**G**. Parameters calculated by the Quantifier for different behaviors elicited by optogenetic

activation of nociceptors. The colors match those for behaviors shown in **A**. Data in **F** and **G** are

from the control group in **A**.

I-**N**. Parameters calculated by the Quantifier for different behaviors elicited by sound stimulation.

The colors match those in **H**.

Probability in **B** and **I** is the fraction of responders (behavioral count > 0) in total animals. n.s., P >

0.05; *, P < 0.05; ***, P < 0.001 (Fisher's exact test for **B** and **I**; unpaired t test for **C**-**G** and **J**-**N**).

Figure 8. The Quantifier captures qualitative and quantitative changes in rodent behaviors.

A. Raster plots showing the frame-wise categorizations by *RatA* of behavior after saline or amphetamine injection. Plots at the left (baseline) show responses to the first exposure to amphetamine across increasing doses (baseline; 0.32, 1.0, 3.2, and 5.6 mg / kg, i.p.); Plots at

the right show responses to increasing doses of amphetamine after repeated amphetamine

treatment (post treatment; 21 days, see methods). Animals in the saline group received

repeated saline throughout. Color bars: behavioral categories (intensities indicate the behavioral

probabilities). X-axis: time (each tick is 10-sec); Y-axis: each row represents an individual rat.

These same colors are used to indicate behavioral categories in panels **B**-**K**.

B-D. The dose response curves of rearing intensity (**B**), locomotion counts (**C**) and head

movement duration (**D**) following the first exposure to amphetamine or saline (baseline). *, $P <$

0.05; **, $P < 0.01$; ***, $P < 0.001$ (unpaired t test between saline and amphetamine groups).

E. Dose response curves for head movement duration at baseline and after repeated

amphetamine exposure (post treatment, i.e., sensitization). *, $P < 0.05$; **, $P < 0.01$; ***, $P <$

0.001 (unpaired t test between baseline and post treatment groups).

F-H. Within subject comparison of rearing intensity (**F**), locomotion distance (**G**) and the head

movement duration (**H**) in response to the first 0.32 mg / kg amphetamine at baseline and after

repeated amphetamine treatment. *, $P < 0.05$ (paired t test).

I-K. Dose response curves of the intensity and counts of body grooming (**I** and **J**) and the

duration of face grooming (**K**) in in rats repeatedly treated with saline.

L. Raster plots showing the frame-wise categorizations by *MouseH* for a younger (32-week) or

an older (80-week) mouse. Color bars: behavioral categories (intensities indicate the behavioral

probabilities). X-axis: time (in second); Y-axis: different random 2-minute time window. Multiple

(>100) 2-minute time windows were randomly selected from a 4-hour recording for each mouse,

and those containing wheel-running behavior (targeted time window) were sorted out. 10

targeted time windows for each mouse were randomly selected for analysis.

M. The running vigor of the two mice in **L**. **, $P < 0.01$ (unpaired t test).

Figure 9. Benchmark comparison between the existing standards and *LabGym*.

A. The workflow of the benchmark comparison between the combination of DeepLabCut and B-SOiD (DLC+B-SOiD) and *LabGym*, which was performed on the same computer that has a 3.6

GHz Intel Core i9-9900K CPU, a 64 GB memory, and an NVIDIA GeForce RTX 2080 GPU).

The setting of Categorizer in *LabGym*: Animation Analyzer: level 1 with input shape 8 X 8 X 1;

Pattern Recognizer: level 2 with input shape 16 X 16 X 3.

B. The total time spent for DLC+B-SOiD and *LabGym* to finish the workflow in **A**. The total

human labor was fixed to 30 minutes for both.

C-D. Comparisons of frame-wise behavior classifications between the computer software and

the ground truth (expert human annotation) in 2 testing videos: baseline (**C**) and post-treatment

(**D**). The raster plots show frame-wise behavior events with the X-axis indicating the time (1

minute). Different colors indicate different behavior types. Videos used here were randomly

selected from those used for Figures 8**A**-8**E**.

E-F. Examples of behavior quantification (locomotion) in baseline (base) and post-treatment

(post) videos by B-SOiD (Ctrl = baseline, Exp = post-treatment; **F**) and *LabGym* (**G**).

G. Comparison of frame-wise behavior classifications between the computer software and the

ground truth (expert human annotation) in generalization to a different testing enclosure. The

raster plots show frame-wise behavior events with the X-axis indicating the time (1 minute). The

colors match those for behaviors shown in **C**.